Atty Dkt No. DETR0101PUSA

## Amendments to the Claims:

This listing of claims will replace all prior versions, and listings, of claims in the application:

## 1-10. (cancelled)

11. (New) Fusion proteins comprising:

regions selected from the group consisting of specific antigen binding regions, microtubule binding regions, and immune response triggering regions.

- 12. (New) The fusion proteins of claim 11 wherein the specific antigen binding regions comprise a ligand or a ligand region.
- 13. (New) The fusions proteins of claim 11 wherein the specific antigen binding regions include a component selected from the group consisting of EGF, FGF, CSF, MGF, IL 15, IL 2, and regions thereof.
- 14. (New) The fusions proteins of claim 11 wherein the specific antigen binding regions include a component selected from the group consisting of gephyrin, Tau, MAP, MID 1, MBP, put MBD or PMBP, FLJ 31424 Fis and regions thereof.
- 15. (New) The fusions proteins of claim 11 wherein the immune response triggering regions include a component selected from the group consisting of Fc of IgG, B 7.1, B 7.2 and regions thereof.
- 16. (New) The fusion proteins of claim 11 further comprising spacer or linker regions.

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- 17. (New) The fusion proteins of claim 16 wherein the spacer regions include polyglycine, polyproline, prolines, or glycines.
- 18. (New) The fusion proteins of claim 11 further comprising at least one component selected from the group consisting of nucleic acid binding regions; polysaccharide binding regions; GFP, fluorescent regions; membrane penetration domains; GST regions; His tags, and regions for carry out of affinity purification.a cleavage site for a protease..
- 19. (New) The fusion proteins of claim 18 wherein the at least one polysaccharide binding region comprises a cellulose binding region of the CipA protein.
- 20. (New) The fusion proteins of claim 18 wherein the GFP, fluorescent regions or the membrane penetration domains include the Gene 3 Protein of the bacteriophage fd, gp 41 or Tat protein of the HIV 1.
- 21. (New) Nucleic Acid and Amino Acid sequences, DNA vectors, amino and nucleic acid, and cloning and expression systems for the fusion proteins of claim 11.

## 22. (New) Fusion proteins comprising:

regions selected from the group consisting of antibody binding regions, and microtubule - binding regions.

- 23. (New) The fusion proteins of claim 22 wherein the antibody binding regions include a component selected from the group consisting of Staphylococcal protein A (SPA), extracellular region of the Fc receptor CD 64, and regions thereof.
- 24. (New) The fusion proteins of claim 22 wherein the microtubule binding regions include a component selected from the group consisting of Tau, MAP, MID 1, MBP, FL J 31424 Fis, and regions thereof.

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- 25. (New) Fusion proteins according to Claim 22 further comprising immune triggering regions.
- 26. (New) The fusion proteins of claim 25 wherein the immune triggering regions include Fc regions of immunoglobulin G or IgG, HLA B 7.1 or HLA B 7.2.
- 27. (New) The fusion proteins of claim 22 further comprising at least one component selected from the group consisting of spacer or linker regions; nucleic acid binding regions; polysaccharide binding regions; GFP, fluorescent regions; membrane penetration domains; GST regions; His tags, and regions for carry out of affinity purification.a cleavage site for a protease.
- 28. (New) The fusion proteins of claim 27 wherein: the polysaccharide binding region comprises a cellulose binding region of the CipA protein;
- the GFP, fluorescent regions or the membrane penetration domains include the Gene 3 Protein of the bacteriophage fd, gp 41 or Tat protein of the HIV 1; and the spacer regions include polyglycine, polyproline, prolines, or glycines.
- 29. (New) Nucleic Acid and Amino Acid sequences, DNA vectors, amino and nucleic acid, and cloning and expression systems for the fusion proteins of claim 22.
- 30. (New) A nucleic acid or amino acid sequence comprises a sequence selected from the group of sequences consisting of SEQ ID NO: 1 12.